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Bioinformatics and Functional Genomics



Jonathan Pevsner

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Pevsner,2003-11-04 Bioinformatics and Functional Genomics provides a broad based introduction to bioinformatics by following three real world examples throughout the book retinol binding protein breast cancer and a calcium binding site C2 The author emphasizes the use of computational tools and databases to study connections between the structure of proteins and genes to function development evolution and disease Readers learn real skills such as how to analyze genes and proteins how to make trees using phylogenetic software how to extract data and how to identify genes and proteins implicated in diseases

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Computational Text Analysis Soumya Raychaudhuri,2006-01-26 This book brings together the two disparate worlds of computational text analysis and biology and presents some of the latest methods and applications to proteomics sequence analysis and gene expression data Modern genomics generates large and comprehensive data sets but their interpretation requires an understanding of a vast number of genes their complex functions and interactions Keeping up with the literature on a single gene is a challenge itself for thousands of genes it is simply impossible Here Soumya Raychaudhuri presents the techniques and algorithms needed to access and utilize the vast scientific text i e methods that automatically read the literature on all the genes Including background chapters on the necessary biology statistics and genomics in addition to practical examples of interpreting many different types of modern experiments this book is ideal for students and researchers in computational biology bioinformatics genomics statistics and computer science

Innovative Methods in Horticultural Crop Improvement Jameel Al-Khayri,Lina M. Alnaddaf,Shri Mohan Jain,Suprasanna Penna,2024-09-30 This book focuses on recent advances in Molecular Tools Nanotechnology and Artificial Intelligence to monitor manage and improve horticultural crops in terms of plant growth nutrient deficiency toxicity diseases abiotic stress soil amendments and agrochemicals entering the surrounding environment The book consists of 11 chapters grouped in 3 parts Part I Nanotechnology and Artificial Intelligence Part II Monitoring Abiotic and Biotic Stress Part III Genetic Engineering and Genomics This book provides an overview of nanotechnology and omics used to improve the productivity of crops and sustainability in the future It also describes the basic structures of six popular artificial intelligence

techniques and their applications in horticultural crop improvement The book presents molecular techniques such as molecular markers genome sequencing genome editing and genetic modification that are considered efficient tools to achieve the goals of plant breeders in horticultural crop improvement programs Chapters are written by globally recognized scientists and subjected to a rigorous review process to ensure quality presentation and scientific precision Each chapter begins with an introduction that covers similar contexts and includes a detailed discussion of the topic accompanied by high quality color images diagrams and relevant details and concludes with recommendations for future study directions in addition to a comprehensive bibliography

Bioinformatics David Edwards,Jason Stajich,David Hansen,2009-09-03

Bioinformatics is a relatively new field of research It evolved from the requirement to process characterize and apply the information being produced by DNA sequencing technology The production of DNA sequence data continues to grow exponentially At the same time improved bioinformatics such as faster DNA sequence search methods have been combined with increasingly powerful computer systems to process this information Methods are being developed for the ever more detailed quantification of gene expression providing an insight into the function of the newly discovered genes while molecular genetic tools provide a link between these genes and heritable traits Genetic tests are now available to determine the likelihood of suffering specific ailments and can predict how plant cultivars may respond to the environment The steps in the translation of the genetic blueprint to the observed phenotype is being increasingly understood through proteome metabolome and phenome analysis all underpinned by advances in bioinformatics Bioinformatics is becoming increasingly central to the study of biology and a day at a computer can often save a year or more in the laboratory The volume is intended for graduate level biology students as well as researchers who wish to gain a better understanding of applied bioinformatics and who wish to use bioinformatics technologies to assist in their research The volume would also be of value to bioinformatics developers particularly those from a computing background who would like to understand the application of computational tools for biological research Each chapter would include a comprehensive introduction giving an overview of the fundamentals aimed at introducing graduate students and researchers from diverse backgrounds to the field and bring them up to date on the current state of knowledge To accommodate the broad range of topics in applied bioinformatics chapters have been grouped into themes gene and genome analysis molecular genetic analysis gene expression analysis protein and proteome analysis metabolome analysis phenome data analysis literature mining and bioinformatics tool development Each chapter and theme provides an introduction to the biology behind the data describes the requirements for data processing and details some of the methods applied to the data to enhance biological understanding

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Food Chemistry Srinivasan Damodaran,Kirk L. Parkin,Owen R. Fennema,2007-09-18 This latest edition of the most internationally respected reference in food chemistry for more than 30 years Fennema s Food Chemistry once again meets and surpasses the standards of quality comprehensive information set by its predecessors This edition introduces new editors

and contributors who are recognized experts in their fields All chapters reflect recent scientific advances and where appropriate have expanded and evolved their focus to provide readers with the current state of the science of chemistry for the food industry The fourth edition presents an entirely new chapter Impact of Biotechnology on Food Supply and Quality which examines the latest research in biotechnology and molecular interactions Two former chapters receive extensive attention in the new edition including Physical and Chemical Interactions of Components in Food Systems formerly Summary Integrative Concepts and Bioactive Substances Nutraceuticals and Toxicants formerly Toxic Substances which highlights bioactive agents and their role in human health and represents the feverish study of the connection between food and health undertaken over the last decade It discusses bioactive substances from both a regulatory and health standpoint Retaining the straightforward organization and detailed accessible style of the original this edition begins with an examination of major food components such as water carbohydrates lipids proteins and enzymes The second section looks at minor food components including vitamins and minerals colorants flavor and additives The final section considers food systems by reviewing basic considerations as well as specific information on the characteristics of milk and the postmortem physiology of edible muscle and postharvest physiology of plant tissues Useful appendices provide keys to the international system of units conversion factors log P values calculation and the Greek alphabet

BIOTECHNOLOGY - Volume XI Horst W. Doelle, J. Stefan Rokem, Marin Berovic, 2009-11-16 This Encyclopedia of Biotechnology is a component of the global Encyclopedia of Life Support Systems EOLSS which is an integrated compendium of twenty one Encyclopedias Biotechnology draws on the pure biological sciences genetics animal cell culture molecular biology microbiology biochemistry embryology cell biology and in many instances is also dependent on knowledge and methods from outside the sphere of biology chemical engineering bioprocess engineering information technology biorobotics This 15 volume set contains several chapters each of size 5000 30000 words with perspectives applications and extensive illustrations It carries state of the art knowledge in the field and is aimed by virtue of the several applications at the following five major target audiences University and College Students Educators Professional Practitioners Research Personnel and Policy Analysts Managers and Decision Makers and NGOs

Biostatistical Genetics and Genetic Epidemiology Robert C. Elston, Jane M. Olson, Lyle Palmer, 2002-04-22 Human Genetics concerns the study of genetic forces in man By studying our genetic make up we are able to understand more about our heritage and evolution Some of the original and most significant research in genetics centred around the study of the genetics of complex diseases genetic epidemiology This is the third in a highly successful series of books based on articles from the Encyclopedia of Biostatistics This volume will be a timely and comprehensive reference for a subject that has seen a recent explosion of interest following the completion of the first draft of the Human Genome Mapping Project The editors have updated the articles from the Human Genetics section of the EoB have adapted other articles to give them a genetic feel and have included a number of newly commissioned articles to ensure the work is comprehensive and provides a self

contained reference Agriculture, Rural Development, Food and Drug Administration, and Related Agencies Appropriations for 2009 United States. Congress. House. Committee on Appropriations. Subcommittee on Agriculture, Rural Development, Food and Drug Administration, and Related Agencies,2008 Fungal Genomics ,2003-08-28 The purpose of this volume is to highlight wide ranging applications of genomics in the area of applied mycology and biotechnology The volume covers a brief overview on fungal genomics meiotic recombination in fungi molecular genetics of circadian rhythms genome sequencing transposable elements mitochondrial genomes ribosome biogenesis pathogenicity genes genetic improvement of yeasts microarrays techniques and applications fungal germplasm and data bases Although it is difficult to develop a comprehensive volume on fungal genomics because of the range and complexity of the emerging knowledge an attempt has been made to bring together pertinent information that will serve the needs of the reader provide a quick reference to material that might otherwise be difficult to locate and furnish a starting point for further study

Bioinformatics Darbeshwar Roy,2009 Issuing in depth coverage of the principles and applications of bioinformatic analyses of nucleic acid and protein sequences databases and data mining discusses dynamic programming algorithms amino acid exchange matrices and pair wise sequence alignment tools in extensive detail

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